

HP95	1	-----MIVADSEC-R-----AFIK-----DYLRF-----PG-GVGDS-----
PA26	1	MAEGENEFVRWDGLCSRDSTTRETALENIROQILRKTEYLRSVKETHRPSDGLSNTESSD
HP95	27	G-----PGEQRES-----RAR-----GPSAFIPVEFVLREG
PA26	61	GLNKLALHLLMLSKRCFKQVREKSEFILKSIQELGIRIPRPLGQGPSRFIPEKFIQVCG
HP95	58	AESELEQH-LGLEALMSSGRVDNIAVMGLHPDYFTSFWXLIHYLLHTDGPLASSWRHYIA
PA26	121	SEDAQMHALFADSAALGRLDNITLVMVFHPQYLESEFLKTHQHYLLQMDGPLPLHYRHYIC
HP95	117	IMAAARHQCSYLVGSHVAEFLQGGDPENLLGLHRAPEKLRKLTSEINKLLAHRPWLITKE
PA26	181	IMAAARHQCSYLVNLHVNDFLHVGGDPKWLINGIENAPQKLNIGENKVL AHRPWLITKE
HP95	177	HIQALEKKTGEHTMSLAELIQAIVLLTHCHSLSSVFGCGILPEGDADGSPAPQAPTPPS-
PA26	241	HIQELLKAEHSHMSLAELVHAVVLLTHYHSLASFTEGCGISPEIHCDGGHTFRPPSVSNY
HP95	236	---EQSSP-PSRD--PIN---NSGGFESARDVEALMERMQLOESLLRDEG-TSQEEMES
PA26	301	CICDIUNGNSVDMPNNSAENVSVSDSFFFEVEALMEKMRQLQE--CRDEEEASQEEMAS
HP95	286	RFELTEKSESLVTPSADILEPSPHPDMLCFVEDPTFGYEDFTRRGAQAPPTFRAQDYTWE
PA26	359	RFELTEKRESMFVFSDD-EEVTPARAVSRHFEDTSYGYKDFSRHGMHVP-TFRVQDYQWE
HP95	346	DHGYSLIGRLYPGGQLDEKFAQAYS LTYNTHAMHSGVDTSVLRRRAINWYIHCVFGIRY
PA26	417	DHGYSLVNRLYPDVGQLDEKFIAYNLTYNTAMHKD VDTSM LRRRAINWYIHCVFGIRY
HP95	406	DDYDYGEVNOILLERNLKVYIKTVACYPEKTT RMYNIEFWRHFRHSEKVVHVNLLLEARMQ
PA26	477	DDYDYGEINOLLDRSFKVYIKTVCTPEKVT RMYDSFWRQFKHSEKVVHVNLLLEARMQ
HP95	466	AALLYALRAITRYMT
PA26	537	AALLYALRAITRYMT

FIGURE 1



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**FIGURE 2**

FIG. 3

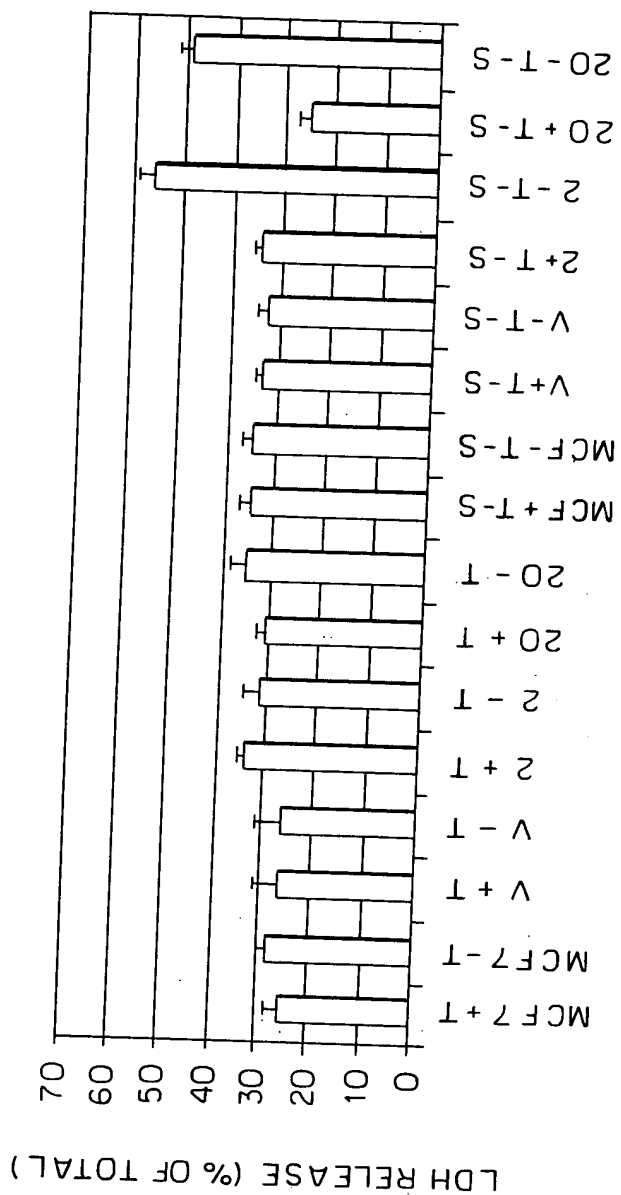


FIG. 4

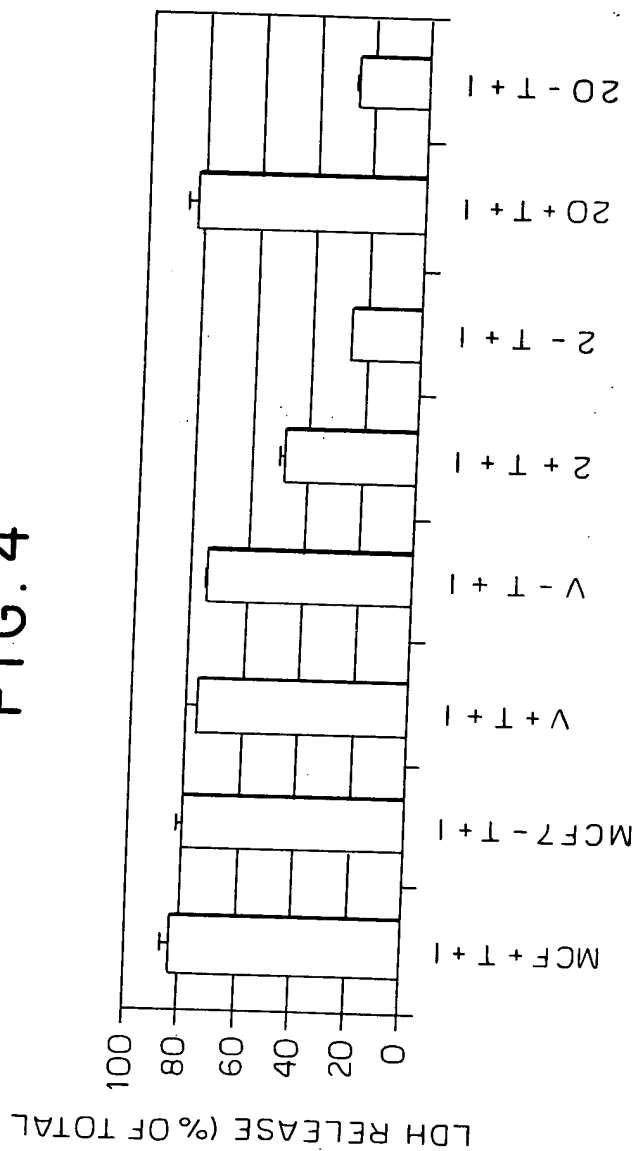


FIG. 5

